

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/720,843
Source: IFW0
Date Processed by STIC: 3/23/05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05



IFW

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/720,843

DATE: 03/23/2005

TIME: 12:14:13

Input Set : N:\DA\pto.da.txt

Output Set: N:\CRF4\03232005\J720843.raw

3 <110> APPLICANT: Solulink
 4 Schwartz, David A.
 6 <120> TITLE OF INVENTION: HYDRAZINE-BASED AND CARBONYL-BASED
 7 BIFUNCTIONAL CROSSLINKING REAGENTS
 10 <130> FILE REFERENCE: 37154-753
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/720,843
 C--> 13 <141> CURRENT FILING DATE: 2003-11-24
 15 <150> PRIOR APPLICATION NUMBER: 60/191,186
 16 <151> PRIOR FILING DATE: 2000-03-22
 18 <160> NUMBER OF SEQ ID NOS: 1
 20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 22 <210> SEQ ID NO: 1
 23 <211> LENGTH: 26
 24 <212> TYPE: DNA
 25 <213> ORGANISM: Artificial Sequence
 27 <220> FEATURE:
 28 <223> OTHER INFORMATION: 25-mer phosphodiester oligonucleotide modified to
 29 incorporate a C6-aminolinker
 W--> 31 <221> NAME/KEY: modified_base
 32 <222> LOCATION: 1
 33 <223> OTHER INFORMATION: N= n-hexylamino (linker)
 W--> 35 <400> 1
 W--> 36 ntttttttag cctaactgat gccatg

**Does Not Comply
Corrected Diskette Needed**

"linker" denotes a small
synthetic single-chain
oligodeoxynucleotide.
26

"N" can only represent
a single nucleotide.

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/720,843

DATE: 03/23/2005
TIME: 12:14:14

Input Set : N:\DA\pto.da.txt
Output Set: N:\CRF4\03232005\J720843.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 1

VERIFICATION SUMMARY

DATE: 03/23/2005

PATENT APPLICATION: US/10/720,843

TIME: 12:14:14

Input Set : N:\DA\pto.da.txt

Output Set: N:\CRF4\03232005\J720843.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:31 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:35 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:36 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0